Example

SSTAR Select samples of monocyte and fibroblast Identify samples of interests **BioLayout** Find SPI1 and its binding motif as active **SSTAR** transcription factor and motif. Find Find potential regulators Enhancer slider enhancers active in monocyte. Confirm the presence of SPI1 at monocyte Confirm gene expression BioGPS plugin by using BioGPS. Inspect alternative TSSs, antisense **ZENBU** Inspect proximal transcription transcription, as well as epigenetic marks **UCSC** and epigenetic status Obtain CAGE peak activity profiles of TET monocyte and fibroblast, perform Perform data analysis BioMart differential analysis, and combine with data files annotation.